
Sequence Listing was accepted with existing errors.

See attached Validation Report.

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217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: Mon May 14 12:25:53 EDT 2007

Validated By CRFValidator v 1.0.2

Application No: 10585880 Version No: 1.1

Input Set:

Output Set:

Started: 2007-05-14 12:25:28.045 **Finished:** 2007-05-14 12:25:29.280

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 235 ms

Total Warnings: 4
Total Errors: 1

No. of SeqIDs Defined: 18

Actual SeqID Count: 18

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
E 224	$<\!220\!>, <\!223\!>$ section required as $<\!213\!>$ has Artificial sequence or Unknown in SEQID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)

SEQUENCE LISTING

<110> CHONNAM NATIONAL UNIVERSITY et al. MUCOSAL VACCINE ADJUVANTS CONTAINING BACTERIAL FLAGELLINS AS <120> AN ACTIVE COMPONENT <130> 095704 <140> 10/585,880 <141> 2006-07-11 <150> KR 10-2004-0001974 <151> 2004-01-12 <160> 18 <170> KopatentIn 1.71 <210> 1 <211> 1131 <212> DNA <213> Vibrio vulnificus <400> atggctatca atgtaaacac taacgtgtca gcaatgaccg cacagcgtta cctaaaccag 60 gccgctgaag gtcaacaaaa atcaatggag cgtttgtctt cgggctataa aatcaatagc gcgaaagatg atgctgcagg tctacaaatt tctaaccgtt tgaactcgca aagccgtggt 180 ctcgacatgg cggttaaaaa tgccaacgat ggtatctcta ttgcacagac tgctgaaggt 240 gcaatgacag agaccaccaa catcctacaa cgtatgcgtg accttgcctt gcaatcgtct 300 aacggttcga actctcgttc tgaacgcgtg gcgattcaag aagaagtgtc agcgttgaac caagaactta accqtatcqc agaqacaacc tcttttqqtq qtaacaaact ccttaacqqt 420 acgtacggtt ctcaatcttt ccaaatcggt gctgactctg gtgaagctgt gatgctttct 480 atgggtaacc ttcgttcaga tacagacgcg atgggcggct tgagctacaa atctgaagaa 540 ggcgtaggcg cagattggcg tgtaagcgac aacactgact tcacgatgtc ttatgtgaat 600 aagcaaggtg aagaaaaaga gatcacagtc aacgccaaag cgggtgacga tcttgaagaa 660 ctggcgactt acatcaacgg tcaaaacgat gatgtgaaag cgtcggtcgg tgaaggcggc 720 780 aaactgcagc tattcgcttc taaccaacgt gtagaaggtg aagtggaatt cggtggtggt ctagcqtctq aqttqaacat tqqtqatqqc accaaaacca atqtqaqcaa cattqatqtc acqacqqttq ctqqctctca aqaaqcaqta qcqatcattq atqqcqcatt qaaatcqqta 900

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960

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Gln Ile Ser Asn Arg Leu Asn Ser Gln Ser Arg Gly Leu Asp Met Ala 50 55 60
Val Lys Asn Ala Asn Asp Gly Ile Ser Ile Ala Gln Thr Ala Glu Gly 65 70 75 80
Ala Met Thr Glu Thr Thr Asn Ile Leu Gln Arg Met Arg Asp Leu Ala 85 90 95
Leu Gln Ser Ser Asn Gly Ser Asn Ser Arg Ser Glu Arg Val Ala Ile 100 105 110
Gln Glu Glu Val Ser Ala Leu Asn Gln Glu Leu Asn Arg Ile Ala Glu 115 120 125
Thr Thr Ser Phe Gly Gly Asn Lys Leu Leu Asn Gly Thr Tyr Gly Ser 130 135 140
Gln Ser Phe Gln Ile Gly Ala Asp Ser Gly Glu Ala Val Met Leu Ser 145 150 155 160
Met Gly Asn Leu Arg Ser Asp Thr Asp Ala Met Gly Gly Leu Ser Tyr 165 170 175
Lys Ser Glu Glu Gly Val Gly Ala Asp Trp Arg Val Ser Asp Asn Thr 180 185 190
Asp Phe Thr Met Ser Tyr Val Asn Lys Gln Gly Glu Glu Lys Glu Ile 195 200 205
Thr Val Asn Ala Lys Ala Gly Asp Asp Leu Glu Glu Leu Ala Thr Tyr 210 215 220
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Lys Leu Gln Leu Phe Ala Ser Asn Gln Arg Val Glu Gly Glu Val Glu
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Phe Gly Gly Leu Ala Ser Glu Leu Asn Ile Gly Asp Gly Thr Lys
260 265 270

Thr Asn Val Ser Asn Ile Asp Val Thr Thr Val Ala Gly Ser Gln Glu 275 280 285

Ala Val Ala Ile Ile Asp Gly Ala Leu Lys Ser Val Asp Ser Glu Arg 290 295 300

Ala Ser Leu Gly Ala Phe Gln Asn Arg Phe Asn His Ala Ile Ser Asn 305 310 315 320

Leu Ser Asn Ile Asn Glu Asn Val Asn Ala Ser Ser Arg Ile Lys 325 330 335

Asp Thr Asp Tyr Ala Lys Glu Thr Thr Gln Met Thr Lys Thr Gln Ile 340 345 350

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<212> PRT

<213> Vibrio vulnificus

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Gln Ile Ser Asn Arg Leu Asn Val Gln Ser Arg Gly Leu Asp Val Ala
50 55 60

Val Arg Asn Ala Asn Asp Gly Ile Ser Ile Ala Gln Thr Ala Glu Gly
65 70 75 80

Ala Met Asn Glu Thr Thr Asn Ile Leu Gln Arg Met Arg Asp Leu Ser 85 90 95

Leu Gln Ser Ala Asn Gly Ser Asn Ser Lys Ser Glu Arg Val Ala Ile 100 105 110

Gln Glu Glu Val Thr Ala Leu Asn Asp Glu Leu Asn Arg Ile Ala Glu 115 120 125

Thr Thr Ser Phe Gly Gly Asn Lys Leu Leu Asn Gly Thr Tyr Gly Thr 130 135 140

Lys Ala Met Gln Ile Gly Ala Asp Asn Gly Glu Ala Val Met Leu Ser 145 150 155 160

Leu Lys Asp Met Arg Ser Asp Asn Val Met Met Gly Gly Val Ser Tyr 165 170 175

Gln Ala Glu Glu Lys Asp Lys Asn Trp Asn Val Ala Ala Gly Asp

180 185 190

Asn Asp Leu Thr Ile Ala Leu Thr Asp Ser Phe Gly Asn Glu Gln Glu 200 Ile Glu Ile Asn Ala Lys Ala Gly Asp Asp Ile Glu Glu Leu Ala Thr 215 220 Tyr Ile Asn Gly Gln Thr Asp Leu Val Lys Ala Ser Val Gly Glu Gly 230 235 Gly Lys Leu Gln Ile Phe Ala Gly Asn Asn Lys Val Gln Gly Glu Ile 245 250 Ala Phe Ser Gly Ser Leu Ala Gly Glu Leu Gly Leu Gly Glu Gly Lys 260 265 Asn Val Thr Val Asp Thr Ile Asp Val Thr Thr Val Gln Gly Ala Gln 280 Glu Ser Val Ala Ile Val Asp Ala Ala Leu Lys Tyr Val Asp Ser His 290 295 300 Arg Ala Glu Leu Gly Ala Phe Gln Asn Arg Phe Asn His Ala Ile Ser 310 315 Asn Leu Asp Asn Ile Asn Glu Asn Val Asn Ala Ser Lys Ser Arg Ile 325 330 Lys Asp Thr Asp Phe Ala Lys Glu Thr Thr Gln Leu Thr Lys Thr Gln 340 345 Ile Leu Ser Gln Ala Ser Ser Ser Ile Leu Ala Gln Ala Lys Gln Ala 355 360 Pro Asn Ser Ala Leu Ser Leu 370 375 <210> 5 <211> 1133 <212> DNA <213> Vibrio vulnificus <400> gtggcgatca ccgttaatac caatgtggca gcacttgtcg cacagcgtca tttgaccagt gcaaccgaca tgctgaatca atccttggag cgtttgtctt cagggaagcg tattaatagt gcaaaagacg atgcggcagg gctgcaaatt tcgaatcgtc ttcagtcgca aatgcgtggt 180 ttagatatcg cggtgcgaaa tgccaatgat ggcatctcca ttatgcagac tgcggaaggg 300 gcaatgaatg aaaccactaa tattctccaa aggatgcgtg atctttcatt gcaatccgcc

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<212> PRT

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Gln Ile Ser Asn Arg Leu Gln Ser Gln Met Arg Gly Leu Asp Ile Ala 50 55 60

Val Arg Asn Ala Asn Asp Gly Ile Ser Ile Met Gln Thr Ala Glu Gly 65 70 75 80

Ala Met Asn Glu Thr Thr Asn Ile Leu Gln Arg Met Arg Asp Leu Ser 85 90 95

Leu Gln Ser Ala Asn Gly Ser Asn Ser Tyr Ala Glu Arg Ile Ala Leu 100 105 110

Gln Glu Glu Met Thr Ala Leu Asn Asp Glu Leu Asn Arg Ile Ala Glu 115 120 125

Thr Thr Ser Phe Gly Gly Arg Lys Leu Leu Asn Gly Ser Phe Gly Ser 135 Ala Ala Phe Gln Ile Gly Ala Ala Ser Gly Glu Ala Val Gln Val Gln 150 155 Leu Lys Ser Met Arg Ser Asp Gly Ile Asp Met Gly Gly Phe Ser Tyr 165 170 Ile Ala Asn Gly Arg Ala Arg Ser Asp Trp Gln Val Lys Glu Gly Ala 185 180 Asn Ala Leu Ser Met Ser Phe Thr Asn Arg Phe Gly Glu Thr Glu Thr 200 Ile Gln Ile Asn Ala Lys Ala Gly Asp Asp Ile Glu Glu Leu Ala Thr 215 220 Tyr Ile Asn Gly Gln Thr Asp Lys Val Thr Ala Ser Val Asn Glu Glu 230 235 Gly Gln Leu Gln Leu Phe Met Ala Gly Glu Glu Thr Ser Gly Thr Leu 245 250 Ser Phe Ser Gly Asp Leu Ala Ser Glu Leu Gly Leu Gln Leu Lys Gly 260 265 Tyr Asp Ala Val Asp Asn Ile Asp Ile Thr Ser Val Gly Ala Gln 275 280 Gln Ala Val Ala Val Leu Asp Thr Ala Met Lys Tyr Val Asp Ser His 290 295 300 Arg Ala Glu Leu Gly Ala Tyr Gln Asn Arg Phe Ser His Ala Ile Asn 310 315 Asn Leu Asp Asn Ile His Glu Asn Leu Ala Thr Ser Asn Ser Arg Ile 330 325 Gln Asp Thr Asp Tyr Ala Lys Glu Thr Thr Arg Met Val Lys Gln Gln Ile Leu Gln Gln Val Ser Thr Ser Ile Leu Ala Gln Ala Lys Lys Gly 360 365 355 Pro Asn Leu Ala Leu Thr Leu 370 375 <210> <211> 1158 <212> <213> Vibrio vulnificus

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ctagatgtgg	cgatgcgtaa	tgccaacgat	ggtatctcta	tcgctcaaac	cgccgaaggg	240
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<213> Vibrio vulnificus

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Ser Ser Gly His Lys Ile Asn Ser Ala Lys Asp Asp Ala Ala Gly Leu 35 40 45

Gln Ile Ser Asn Arg Leu Thr Ala Gln Ser Arg Gly Leu Asp Val Ala 50 55 60

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His	Glu	Glu 115	Ala	Ser	Ala	Leu	Gln 120	Asp	Glu	Ile	Asn	Arg 125	Ile	Ala	Glu	
Thr	Thr 130	Ser	Phe	Gly	Gly	Arg 135	Arg	Leu	Leu	Asn	Gly 140	Thr	Phe	Gly	Asp	
Ala 145	Ala	Phe	Gln	Ile	Gly 150	Ser	Asn	Ser	Gly	Glu 155	Ala	Met	Ile	Met	Gly 160	
Leu	Thr	Ser	Ile	Arg 165	Ala	Asp	Asp	Phe	Arg 170	Met	Gly	Gly	Thr	Thr 175	Phe	
Gln	Ser	Glu	Asn 180	Gly	Lys	Asn	Lys	Asp 185	Trp	Glu	Val	Ser	Ala 190	Asp	Asn	
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Ile 225	Glu	Glu	Leu	Ala	Thr 230	Tyr	Ile	Asn	Gly	Gln 235	Ser	Asp	Tyr	Ile	Asn 240	
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